SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT:

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(ii) TITLE OF INVENTION: Polyketides and their synthesis

(iii) NUMBER OF SEQUENCES: 93

(iv) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk

(B) COMPUTER: IBM PC compatible

(C) OPERATING/SYSTEM: PC-DOS/MS-DOS

(D) SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)

(v) CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/GB97/01819

(vi) PRIOR APPLICATION DATA:

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(vi) PRIOR APPLICATION DATA:

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(B) FILING DATE: 28-MAY-1997

(vi) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: US 60/024188

(B) FILING DATE: 19-AUG-1996

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 56 base pairs

(B) / TYPE: nucleic acid

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
ACGC	GUACUA GUCCGATTAA TTAAGGAGGA CCATCATGGC GGACCTGTCA AAGCTC	56
(2)	INFORMATION FOR SEQ ID NO:2:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 63 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
AUGG	GAGAUCU CUCCGCTAGC GGTTCGCCGG GCGCCGCTTC GTTGGTCCGC GCGCGGGTTT	60
CCC		63
(2)	INFORMATION FOR SEQ ID NO:3:	
•	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
ACTA	AGTCCAC TGCCTCTCGG TAAAATCCAG C	31
(2)	INFORMATION FOR SEQ ID NO:4:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
СТТА	AAGAGGG GCTCCACCGC GTTCACGGAC	3 (
(2)	INFORMATION FOR SEQ ID NO:5:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

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	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:	
ACA'	TTCTC	TA CGCCTAAGTG TTCCCCTCCC TGCCTC	36
(2)	INFO	RMATION FOR SEQ ID NO:6:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 53 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:6:	
GTG	ATGTA'	TG CTCATATGTG TCCTCCTTAA TTAATCGATG CGTTCGTCCG GTG	53
(2)	INFO	RMATION FOR SEQ ID NO:7:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:7:	
TGA	ACACC	AA GCTTGCCAGA GAGCGACGAC TTCCCC	36
(2)	INFO	RMATION FOR SEQ ID NO:8:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:8:	
GAC	AGATT	GC ATGCCCTTCG AGGAGTGCCC GCCCGG	36
(2)	INFO	RMATION FOR SEQ ID NO:9:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:9:	

GACAGATTCT CGAGCCTTCG AGGAGTGCCC GCCCGG	36
(2) INFORMATION FOR SEQ ID NO:10:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
TAAGGAGGAC ACATATGCA	19
(2) INFORMATION FOR SEQ ID NO:11:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
TAATTCCTCC TGTGTAT	17
(2) INFORMATION FOR SEQ ID NO:12:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
CCCATATGGC GGACCTGTCA AAGC	24
(2) INFORMATION FOR SEQ ID NO:13:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
ATTGCGCGCC CTGGCCCGGG AA	22
(2) INFORMATION FOR SEQ ID NO:14:	

	(A) LENGTH: 22 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:	
AAT	TCATAGT CTAGAAGCTT AT	22
(2)	INFORMATION FOR SEQ ID NO:15:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
CGA'	TAAGCTT CTAGACTATG	20
(2)	INFORMATION FOR SEQ ID NO:16:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
СТС	GTCGGTG GCTTTGCG	18
(2)	INFORMATION FOR SEQ ID NO:17:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
CCC	GGGAAAA ACGAAGACTA GTGGCGCGGA CGGCCG	36
(2)	INFORMATION FOR SEQ ID NO:18:	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 18 base pairs(B) TYPE: nucleic acid	

(i) SEQUENCE CHARACTERISTICS:

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
CACG	CGCAGC GCGGCGGA	18
(2)	INFORMATION FOR SEQ ID NO:19:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
CGAA	CCGCTA GCGGTCGTCG CGATGGCCT	29
(2)	INFORMATION FOR SEQ ID NO:20:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
GTGG	CCCGGC CGTCCGCGC ACTAGTCTTC GTTTTT	36
(2)	INFORMATION FOR SEQ ID NO:21:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
AACA	AGCTAGC GGTTCGTCCG CCGCTGCCGT GCC	33
(2)	INFORMATION FOR SEQ ID NO:22:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 48 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear 	

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:	
AATTCACATC ACCATCACCA TCACTAGTAG GAGGTCTGGC CATCTAGA	48
(2) INFORMATION FOR SEQ ID NO:23:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 46 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:	
CGCTTCTAGA TGGCCAGACC TCCTACTAGT GATGGTGATG GTGATG	46
(2) INFORMATION FOR SEQ ID NO:24:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:	
GCAGGGATAT CGCACGTTCC TGG	23
(2) INFORMATION FOR SEQ ID NO:25:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:	
CGCCGAGATC TGCGAAGGCC TGGTCGGCGG G	31
(2) INFORMATION FOR SEQ ID NO:26:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:	
ATGAATTCCC TCCGCCCAGC CAG	23

 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:	
ACAGATCTCG GCTTCGACTC GCTGACCG	28
(2) INFORMATION FOR SEQ ID NO:28:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 55 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:	
ACGCGUACUA GUCAGATCTG GGCATCAATT CGCTGACCGC GGTGGAACTG CGCAA	55
(2) INFORMATION FOR SEQ ID NO:29:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 54 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:	
AUGGAGAUCU CUCAGATCTT GAATGCGGCG GCTGCGGGGA TGGTGCTGGC GTCA	54
(2) INFORMATION FOR SEQ ID NO:30:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 45 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:	
GCACCCACGA CGCCACCACC ACATATGCCC TGCACCCTGC CCTCC	45
(2) INFORMATION FOR SEQ ID NO:31:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 57 base pairs	

(2) INFORMATION FOR SEQ ID NO:27:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:	
ACGCGUACUA GUCCGATTAA TTAAGGAGGA CCATCAATGG CGGACCTGTC AAAGCTC	57
(2) INFORMATION FOR SEQ ID NO:32:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:	
AUGGAGAUCU CUCCGCTAGC GATTGTGGGT ATGGCG	36
(2) INFORMATION FOR SEQ ID NO:33:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 47 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33: ACGCGUACUA GUCCATGCAT CTGCAGCACG GCGGCCTCAT CACCGGA	47
(2) INFORMATION FOR SEQ ID NO:34:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 41 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:	
ACGCGUACUA GUCCATGCAT TCCCGGAGCG GCGATCTGTG G	41
(2) INFORMATION FOR SEQ ID NO:35:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 53 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:	
AUGGAGAUCU CUCCCGCGGC CGCGCTGTCA CGCACCAGCT TCAGCAGTGC GTC	53
(2) INFORMATION FOR SEQ ID NO:36:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 41 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:	
ACGCGUACUA GUCCGCGGCC GCGATCCTCG GGCATTCCAG C	41
(2) INFORMATION FOR SEQ ID NO:37:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	,
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:	
AUGGAGAUCU CUAAGCATTG GTAACTGTC	29
(2) INFORMATION FOR SEQ ID NO:38:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 45 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:	
ACGCGUACUA GUATCTAGAC CATGCATGTT TGACAGCTTA TCATC	45
(2) INFORMATION FOR SEQ ID NO:39:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 45 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:	
AUGGAGAUGU CUATCTAGAC CATGCATGCC GCCGGCTTCC ATTCA	45

	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:	
GAGC	AGTCGT TCCGAGATCT CGGCTTCGAT TCA	33
(2)	INFORMATION FOR SEQ ID NO:41:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
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	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:	
GGGA	GGAGAT CAGATCCCAG AAGT	24
(2)	INFORMATION FOR SEQ ID NO:42:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:	
GAGG	GAGTCG AACCGAGATC TCGGAACGCG CGG	33
(2)	INFORMATION FOR SEQ ID NO:43:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:	
GGGG	GATCCT GGGGTCGGCC GGGCAGGGCA A	31
(2)	INFORMATION FOR SEQ ID NO:44:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs	

(2) INFORMATION FOR SEQ ID NO:40:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:	
GTCTCAAGCT TCGGCATCAG CGGCACCAA	29
(2) INFORMATION FOR SEQ ID NO:45:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:	
CGTGCGATAT CCCTGCTCGG CGAGCGCA	28
(2) INFORMATION FOR SEQ ID NO:46:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 32 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:	
CATGGCCTGC AGGCTGCCCG GGGAGGTCGA CT	32
(2) INFORMATION FOR SEQ ID NO:47:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 35 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:	
CCCGAAGCTT GACACACCTG CCCGGCGCAC CCCGT	35
(2) INFORMATION FOR SEQ ID NO:48:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

(X1) SEQUENCE DESCRIPTION: SEQ ID NO:48:	
GCGCGCCAAT TGCGTGCACA TCTCGAT	27
(2) INFORMATION FOR SEQ ID NO:49:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 37 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:	
CCTGCAGGCC ATCGCGACGA CCGCGACCGG TTCGCCG	37
(2) INFORMATION FOR SEQ ID NO:50:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 32 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:	. 32
GATGGCCTGC AGGCTGCCCG GCGGTGTGAG CA	32
(2) INFORMATION FOR SEQ ID NO:51:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	,
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:51:	
GCCGAAGCTT GAGACCCCCG CCCGGCGCGC TCGC	34
(2) INFORMATION FOR SEQ ID NO:52:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:52:	
TGGCTTCGCT GGCGGACACG CTCAG	25

 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 32 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:53:	
CCTGCAGGCC ATGCCGACGA TCGCGATCGG CT	32
(2) INFORMATION FOR SEQ ID NO:54:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:54:	
GTCAAGCTTC GGGGTGAGCG GGACGAA	27
(2) INFORMATION FOR SEQ ID NO:55:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:	
GCGTCCGGAC GTGGCTCCAG CA	22
(2) INFORMATION FOR SEQ ID NO:56:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:	
GGAGTACTGC GAGGGCGTGG GCAT	24
(2) INFORMATION FOR SEQ ID NO:57:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs	

(2) INFORMATION FOR SEQ ID NO:53:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:57:	
	0.6
CACCTAGGAC CGCTTCCCAG TCGACC	26
(2) INFORMATION FOR SEQ ID NO:58:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 37 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:58:	
TACCTAGGCC GGGCCGGACT GGTCGACCTG CCGGGTT	37
(2) INFORMATION FOR SEQ ID NO:59:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:59:	
ATCCTCAGGC TCTCCGTCTC CGGTTCTCC	29
(2) INFORMATION FOR SEQ ID NO:60:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:60:	
TACCTGAGGG ACCGGCTAGC GGGTCTGCCG CGTG	34
(2) INFORMATION FOR SEQ ID NO:61:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:61:	
CTTCTAGACT ATGAATTCCC TCCGCCCAGC	30
(2) INFORMATION FOR SEQ ID NO:62:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:62:	
CGCCTAGGCA CCACCACAC CCGGGTACTG GACC	34
(2) INFORMATION FOR SEQ ID NO:63:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:	
TAGCTAGCCG GGCGCTCAGG GGCTGCGAGC CGACCT	36
(2) INFORMATION FOR SEQ ID NO:64:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:	
CCTAGGCACC ACCACGGCCC GGGTGCTGGA CCTT	34
(2) INFORMATION FOR SEQ ID NO:65:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:	
CCTCAGGCTG TCACCGGTAG AGGCGGCCCT	30

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(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 32 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:	
ATCCTAGGAC CGCTTCCCAG TCGACCGCGA CA	32
(2) INFORMATION FOR SEQ ID NO:67:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:67:	
GCCACTAGTG TGGCGTGGGG GCTGTGGG	28
(2) INFORMATION FOR SEQ ID NO:68:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:68:	
TGAATTCCCT CCGCCCAGCC AGGCGTCGAT	30
(2) INFORMATION FOR SEQ ID NO:69:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:69:	
CCTGGAGTAC TGCGAGGGCG TG	22
(2) INFORMATION FOR SEQ ID NO:70:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 32 base pairs	

(2) INFORMATION FOR SEQ ID NO:66:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:70:	
CTGACTAGTG GCGGTGACGT GGGCGGGGGA AA	32
(2) INFORMATION FOR SEQ ID NO:71:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 35 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:71:	
CCCCTGCAGC CGGACCGCAC CACCCCTCGT GACGA	35
(2) INFORMATION FOR SEQ ID NO:72:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:72:	
GGCGGAACGT CTTCCCGGCG GCACCT	26
(2) INFORMATION FOR SEQ ID NO:73:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:73:	
CCCCTGCAGC CAGTACCGCT GGGGCTCGAA	30
(2) INFORMATION FOR SEQ ID NO:74:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:74:	
CATGCTCGAG CTCTCCTGGG AAGT	24
(2) INFORMATION FOR SEQ ID NO:75:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:75:	
CAACCCTGGC CAGGGAAGAC GAAGACGG	28
(2) INFORMATION FOR SEQ ID NO:76:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:76:	
TTCCCTGGCC AGGGGTCGCA GCGTG	25
(2) INFORMATION FOR SEQ ID NO:77:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:77:	
CACCTAGGAC CGCGGACCAC TCGAC	25
(2) INFORMATION FOR SEQ ID NO:78:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:78:	
TOCCCACCCA CTCCCTCCAC CTACCCA	27

 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:79:	
GCCGACAGCG AGTCGACGCC GAGTT	25
(2) INFORMATION FOR SEQ ID NO:80:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:80:	
CTGGCCAGGG CGCGCAATGG CCGAGCAT	28
(2) INFORMATION FOR SEQ ID NO:81:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:81:	
CCCTAGGAGT CGCCGGCAGT CCAGCGCGGC GCCC	34
(2) INFORMATION FOR SEQ ID NO:82:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:82:	
AATTCATAAC TAGTAGGAGG TCTGGCCATC TAGA	3
(2) INFORMATION FOR SEQ ID NO:83:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs	

(2) INFORMATION FOR SEQ ID NO:79:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:83:	
TCGAAGATCT ACCGGTCTGG AGGATGATCA ATAC	34
(2) INFORMATION FOR SEQ ID NO:84:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:84:	
GGCGGGTCCG GAGGTGTTCA CCGAGTT	27
(2) INFORMATION FOR SEQ ID NO:85:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:85: ACCTTGGCCA GGGAAGACGA ACACTGA	27
(2) INFORMATION FOR SEQ ID NO:86:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i) GROVENOR PROGRESSMENT GROVEN GROVEN AND OCC	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:86:	36
TCCTAGGCCG GGCCGGACTG GTCGACCTGC CGGGTT	30
(2) INFORMATION FOR SEQ ID NO:87: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:87:	·
AAACACCGCG ACCTGGTCCT CCGAGC	26
(2) INFORMATION FOR SEQ ID NO:88:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:88:	
TTTGCTAGCG ATCGTCGGCA TGGCGTGCCG GTT	33
(2) INFORMATION FOR SEQ ID NO:89:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:89:	
CCCACGAGAT CTCCAGCATG ATCC	24
(2) INFORMATION FOR SEQ ID NO:90:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	·
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:90:	
GGCGGCATGC GGCGGTTCCT	20
(2) INFORMATION FOR SEQ ID NO:91:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 50 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:91:	
AAGCTAGCGG TTCGCCGGGC GCCGCTTCGT TGGTCCGCGC GCGGGTTAAC	50



(2) INFORMATION FOR SEQ ID NO:92:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 40 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:92:

AACCCGCGCG CGGACCAACG AAGCGGCGCC CGGCGAACCG

40

- (2) INFORMATION FOR SEQ ID NO:93:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 44 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear